L3 STRUCTURE UPLOADED

=> s 13 sss full FULL SEARCH INITIATED 17:21:38 FILE 'REGISTRY' FULL SCREEN SEARCH COMPLETED - 54203 TO ITERATE

100.0% PROCESSED 54203 ITERATIONS SEARCH TIME: 00.00.01

3441 ANSWERS

L4 3441 SEA SSS FUL L3

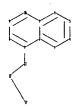
=> s rpr/sqsp

L6 251255 RPR/SQSP

=> s, 14 and 16

1571 L4 31174 L6

L7 18 L4 AND L6



```
chain nodes :
11  12  17
ring nodes :
1  2  3  4  5  6  7  8  9  10
chain bonds :
7-11  11-12  12-17
ring bonds :
1-2  1-6  2-3  2-7  3-4  3-10  4-5  5-6  7-8  8-9  9-10
exact/norm bonds :
7-11  11-12  12-17
normalized bonds :
1-2  1-6  2-3  2-7  3-4  3-10  4-5  5-6  7-8  8-9  9-10
```

G1:CH,NH

G2:0,S

Match level:
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:CLASS 12:CLASS 17:CLASS

(FILE 'HOME' ENTERED AT 17:14:40 ON 21 JUN 2007)	(FILE	'HOME'	ENTERED	AT	17:14:40	ON	21	JUN	2007)
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- FILE 'REGISTRY' ENTERED AT 17:14:55 ON 21 JUN 2007
- L1 STRUCTURE UPLOADED
- L2 0 S L1 SSS FULL
 - FILE 'STNGUIDE' ENTERED AT 17:15:33 ON 21 JUN 2007
- FILE 'REGISTRY' ENTERED AT 17:21:24 ON 21 JUN 2007
- L3 STRUCTURE UPLOADED
- L4 3441 S L3 SSS FULL
- FILE 'HCAPLUS' ENTERED AT 17:21:42 ON 21 JUN 2007 L5 125 S L4 AND PEPTIDE?
- FILE 'REGISTRY' ENTERED AT 17:22:10 ON 21 JUN 2007 L6 251255 S RPR/SQSP
- FILE 'HCAPLUS' ENTERED AT 17:22:26 ON 21 JUN 2007 L7 18 S L4 AND L6
 - FILE 'STNGUIDE' ENTERED AT 17:23:49 ON 21 JUN 2007